```
RESULT 1
US-11-055-822-989
; Sequence 989, Application US/11055822
; Publication No. US20050260707A1
; GENERAL INFORMATION:
   APPLICANT: Pompejus, Markus
   APPLICANT:
             Kroger, Burkhard
   APPLICANT: Schroder, Hartwig
   APPLICANT: Zelder, Oskar
   APPLICANT: Haberhauer, Gregor
   TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING
   TITLE OF INVENTION: METABOLIC PATHWAY PROTEINS
   FILE REFERENCE: BGI-121CPCN
   CURRENT APPLICATION NUMBER: US/11/055,822
   CURRENT FILING DATE: 2005-02-11
   PRIOR APPLICATION NUMBER: 09/606,740
   PRIOR FILING DATE: 2000-06-23
   PRIOR APPLICATION NUMBER: 60/141,031
   PRIOR FILING DATE: 1999-06-25
   PRIOR APPLICATION NUMBER: 60/142,101
   PRIOR FILING DATE: 1999-07-02
   PRIOR APPLICATION NUMBER: 60/148,613
   PRIOR FILING DATE: 1999-08-12
   PRIOR APPLICATION NUMBER: 60/187,970
   PRIOR FILING DATE: 2000-03-09
   PRIOR APPLICATION NUMBER: DE 19930476.9
   PRIOR FILING DATE: 1999-07-01
   PRIOR APPLICATION NUMBER: DE 19931415.2
   PRIOR FILING DATE: 1999-07-08
   PRIOR APPLICATION NUMBER: DE 19931418.7
   PRIOR FILING DATE: 1999-07-08
   PRIOR APPLICATION NUMBER: DE 19931419.5
   PRIOR FILING DATE: 1999-07-08
   PRIOR APPLICATION NUMBER: DE 19931420.9
   PRIOR FILING DATE: 1999-07-08
   Remaining Prior Application data removed - See File Wrapper or PALM.
  NUMBER OF SEQ ID NOS: 1158
; SEQ ID NO 989
   LENGTH: 798
    TYPE: DNA
    ORGANISM: Corynebacterium glutamicum
    FEATURE:
   NAME/KEY: CDS
   LOCATION: (101)..(775)
   OTHER INFORMATION: RXA00105
US-11-055-822-989
  Query Match
                        11.6%; Score 102; DB 12; Length 798;
  Best Local Similarity
                        100.0%; Pred. No. 4.6e-20;
 Matches 102; Conservative 0; Mismatches
                                                0; Indels
                                                              0; Gaps
                                                                         0;
         778 CATCTTGATCTGTCGCCCTGAACTTGCAGATGAACTTCTCGAGATGTGCGCGAAGTTCTA 837
Qу
             1 CATCTTGATCTGCCCCTGAACTTGCAGATGAACTTCTCGAGATGTGCGCGAAGTTCTA 60
         838 CGAGGAGAATGGAACTTACTAACGCTGTTATGATGACGGCAT 879
Qу
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